

Raw Sequence Listing before editing,
for reference only

BEST AVAILABLE COPY

CRF Errors Edited by the STIC Systems Branch

Serial Number:

10516406

CRF Edit Date:

2-1-06

Edited by:

RL

___ Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line

___ Corrected the SEQ ID NO. Sequence numbers edited were:

___ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

☒ Deleted: ☒ invalid beginning/end-of-file text ; ___ page numbers

___ Inserted mandatory headings/numeric identifiers, specifically:

___ Moved responses to same line as heading/numeric identifier, specifically:

___ Other:

Revised 09/09/2003

BEST AVAILABLE COPY



PCT

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/516,406

DATE: 01/26/2006

TIME: 08:44:51

Input Set : A:\Centre sequence listing [1].txt

Output Set: N:\CRF4\01262006\J516406.raw

3 <110> APPLICANT: Centre for Addiction and Mental Health
 5 <120> TITLE OF INVENTION: Detection of Epigenetic Abnormalities and Diagnostic Method
 Based Thereon
 7 <130> FILE REFERENCE: 04340-PCT-PA(0148.0041)
 9 <140> CURRENT APPLICATION NUMBER: 10/516,406
 10 <141> CURRENT FILING DATE: 2004-12-01
 12 <150> PRIOR APPLICATION NUMBER: PCT/CA03/00820
 13 <151> PRIOR FILING DATE: 2003-06-06
 15 <150> PRIOR APPLICATION NUMBER: 60/386,818
 16 <151> PRIOR FILING DATE: 2002-06-06
 18 <160> NUMBER OF SEQ ID NOS: 263
 20 <170> SOFTWARE: PatentIn version 3.1

Does Not Comply
 Corrected Diskette Needed
 (Pg. 1)

ERRORED SEQUENCES

6591 <210> SEQ ID NO: 263
 6592 <211> LENGTH: 566
 6593 <212> TYPE: DNA
 6594 <213> ORGANISM: Homo sapiens
 6596 <220> FEATURE:
 6597 <221> NAME/KEY: misc_feature
 6598 <223> OTHER INFORMATION: Alu sequence cloned from BD43-14 (191, 100, 16q24.2)
 withM13R (se
 6599 e Figure 3)
 6602 <400> SEQUENCE: 263
 6603 ggagggtggt tgcacaatct cagctcacca caaccttttc ctgctggggt caagtgatta 60
 6605 tcctgcctca acctcccgcac tagctgggat tacaggcatg caccaccatg cctggctaata 120
 6607 tttgtatttt tagcagagac agtgtttctc catgttggtg aggctggtct caaactcccg 180
 6609 acctcaggtg atccgcctgc ctcagcctcc caaactgctg ggagtacagg caagccgaat 240
 6611 tctgcagata tccatcacac tggcggccgc tcgagcatgc atctagaggg cccaattcgc 300
 6613 cctatagtga gtcgtattac aattcactgg ccgtcgtttt acaacgtcgt gactgggaaa 360
 6615 accctggcgt taccacaactt aatcgccttg cagcacatcc cccttcgcc agctggcgta 420
 6617 atagcgaaga ggcccgcacc gatcgccctt cccaacagtt gcgcagcctg aatggcgaat 480
 6619 ggacgcgccc tgtaacggcg cattaagcgc ggcggggtgt gtggttacgc gcagcgtgac 540
 6621 cgctacactt gccagcgccc tagcgc 566
 E--> 6626 1

deleted

RAW SEQUENCE LISTING ERROR SUMMARY

DATE: 01/26/2006

PATENT APPLICATION: US/10/516,406

TIME: 08:44:52

Input Set : A:\Centre sequence listing [1].txt

Output Set: N:\CRF4\01262006\J516406.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 5

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:2,3,4,5

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/516,406

DATE: 01/26/2006

TIME: 08:44:52

Input Set : A:\Centre sequence listing [1].txt

Output Set: N:\CRF4\01262006\J516406.raw

L:885 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:540
 L:1001 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45 after pos.:0
 L:1123 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49 after pos.:0
 L:1273 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52 after pos.:420
 L:1275 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52 after pos.:480
 L:1277 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52 after pos.:540
 L:1529 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60 after pos.:360
 L:5543 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:227 after pos.:0
 L:6626 M:254 E: No. of Bases conflict, this line has no nucleotides. /